

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/024,607

DATE: 01/15/2002
TIME: 18:09:09

Input Set : A:\B08017231.txt
Output Set: N:\CRF3\01152002\J024607.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Lee, Richard T.
5 <120> TITLE OF INVENTION: CARDIOVASCULAR DISEASE DIAGNOSTIC AND THERAPEUTIC TARGETS
7 <130> FILE REFERENCE: B0801/7231/ERP/KA
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/024,607
C--> 9 <141> CURRENT FILING DATE: 2001-11-08
9 <150> PRIOR APPLICATION NUMBER: US 60/247,457
10 <151> PRIOR FILING DATE: 2000-11-09
12 <160> NUMBER OF SEQ ID NOS: 33
14 <170> SOFTWARE: FastSEQ for Windows Version 3.0

ERRORED SEQUENCES

488 <210> SEQ ID NO: 9
489 <211> LENGTH: 5028
490 <212> TYPE: DNA
491 <213> ORGANISM: Rattus norvegicus
493 <220> FEATURE:
494 <221> NAME/KEY: mRNA
495 <222> LOCATION: (1)...(5028)
496 <223> OTHER INFORMATION: Lot-1
498 <400> SEQUENCE: 9
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501 ggaccgacgg tgcgcggccga atccggctcg gataggtctg gttggagctt gtgcctgtt 180
502 gcttggcgttggtttcc tgcttgatttgc acgggtgcc attggcttcg tatttggaa 240
E--> 503
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505 tggtgctgcc ttttcagagt caggctgttag cgactccccg cttcgttccc ggctgggctt 480
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512 gaaataccac accatgtctgg gctacaagag gcacatggcc ctgcattcgg ccaggcagg 900
513 cgatctcacc tgcggcgttcc gcacccttgc gctggggagc accgaggatcc tgctggacca 960
514 cctcaagtct cacgcggaaag aaaaggccca ccacgcggcc agggagaaga aacaccatgt 1020
515 cgaccactgc gagagatgtt tctacacccg gaaggatgtg cgtcgcacc tgggtgtcca 1080
516 cacaggatgc aaggacttcc tgcgtcgtt ctgcggccag agattttggc gcaaagacca 1140
517 cctcaactgtt cacaccaaga agacccactc ccaggagctg atgcaagaga gcctgcacg 1200
518 aggagaatac cagggcgggtt accaaccat tgcgcctccg ttccagatca aggtgtatcc 1260
519 catgcctcccttccatgttag aaatgcggcccgagagcggtt cttgtatgggg gcttgcctcc 1320
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523 cccgttcgttccatgtccaaatgc tgcgttccatgttag aaatgcggcccgagagcggtt cttgtatgggg gcttgcctcc 1560

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524	tgaggaattt cctctgcaag agtgcagtc gcctgtcaag ttca	gactgtcagt gctttgagat	1620			
525	ggctaaggaa gggtttggga aagtcaccct gcccaaagag ctgctggtag	atgctgtaaa	1680			
526	tatagccatt cctgctc tggagatttc ctctcttgc	gggttctggc agctgcccc	1740			
527	tcctcctccc cagaatggct tcatgaatgg caccatccct	gtggggcccg gggagccgct	1800			
528	gccccatagg ataacttgtc tggcacagca gcagccacca	cctctgtac ctccgccc	1860			
529	gcccgtccg ctgcagagc cgctgccaca gccacagctg	ccgcccacagt ttca	1920			
530	gtccagccc cagccccaga tgcagccca gatgcagctg	cagecttac agtgcagct	1980			
531	gccccagctg ctgccccagc tgcagccca gcctgagcca	gagccagagc cagaggaaga	2040			
532	agaggaagaa gaagaagaga tagaagaaga agaagagatc	gaagaagaag aagaagccga	2100			
533	accagaagca gaagaagaag aggaggcaga agacaa	gaggcagagg aagaggaaga	2160			
534	agagccacag ccagaagaag cccaaatagc aatagagcgt	gtgaatatgg gccagcccc	2220			
535	gctacccccc acccctcatg tttcacagc tggcaccaac	actgctatcc tgcccccattt	2280			
536	ccaccacgcg ttca	ttccataaa ttgggtttt aagagggtgc ttctttctg	gaagatgttt	2340		
537	caaaccacag ttccagttcc agacatca	gtagttga agagaagcgt tgaaaaaaca	2400			
538	ggaatgggt ttctagctt tgc	ccatgag tagattgaga aaaagaactc tcttaactgc	2460			
539	atgcactgtg ccaatacata tatata	tata tatata tatata tatata	2520			
540	tatata	tatatac atccttagta ttcatgc	tttgc	2580		
541	ctgaaatcgaa ctgcaagtag	tatcatatta ttaccctgat	attgttagtc tcatattatt	2640		
542	agccttgat tattctcata	taatcaaaac caagatccaa aacatgagct	gctaatttgc	2700		
543	aaatatcgtg ttgagtttgc	ggcgtag tgatgttagc tgcgtat	tgatgttagca	2760		
544	ctgccttaga agggcacgag	ggccaagg	ggcttctccc acttggaa	2820		
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546	gtctaccctc aggattggat	gtatgttagt cgcttctt	cttagctc	2940		
547	gtccttctgg tagctactg	tgtgtgtctg tgcgtat	tataggaa	3000		
548	agctgtcggt gtcttgc	tttcaaaaatgt aattttatg	tattttca	3060		
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552	ttgagaaacg tactgaaatt	aaaaaaa aagcttaat	acattagata agaattttct	3300		
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561	taccctaacc cttgaagact	gtatttgc	tttgcgt	tatttcccag	3900	
562	ctaaccctt	gaaactctgt	tatttccattt	tttgcgt	tttgcgt	3960
563	taaccctt	aaagactccat	tttgcgt	tccccc	4020	
564	ttggaggctt	tgtat	tttgcgt	tttgcgt	4080	
565	aagactctgt atttcat	tttgcgt	tttgcgt	tttgcgt	4140	
566	gactctgtat	tttgcgt	tttgcgt	tttgcgt	4200	
567	actctgtatt	tcatttttgcgt	tttgcgt	tttgcgt	4260	
568	tgtat	tttgcgt	tttgcgt	tttgcgt	4320	
569	tatttgcgtt	tttgcgt	tttgcgt	tttgcgt	4380	
570	tttgcgtt	tttgcgt	tttgcgt	tttgcgt	4440	
571	tttgcgtt	tttgcgt	tttgcgt	tttgcgt	4500	
572	tttgcgtt	tttgcgt	tttgcgt	tttgcgt	4560	

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1766	ttatttgcaa	gtcaatgaac	caaacggcg	accaggctcg	ttgtccgcg	ttgtgaaagc	2100
1767	aaggttat	ttatcgccc	ttgccccact	gaacaattc	actgaaaagg	aagagtccca	2160
1768	gccgtgtgt	tgcgcgtgg	gccatacggg	acgtgcagct	acgtgccac	ctccagaacg	2220
1769	actttat	caaagcgatt	accacgttat	ctatttgtt	tcctttcca	gcaagagcag	2280
1770	ccttact	ccctcaaatt	tcttaattac	aaacccgttt	gcttctaaat	caaccccaa	2340
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1773	gatcgctg	tttggttaca	gagcagagac	gcctcagctc	ggatgcaaa	gctaccaaga	2520
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1807	acttcc	gaga	ttgcacac	aagcac	gcgc	caccc	4560
1808	gcagc	ccccgg	tctggc	gctc	cggc	ggcc	4620
1809	gcggc	ggc	ggcaacat	ccgc	ggcccac	cccg	4680
1810	tgctgc	caatgt	gacactg	tcatgc	ggaagt	ttt	4740
1811	tgataga	aaat	gggttgg	cgcatca	actctgg	ggcaaaac	4800
1812	at	tgacgg	ttcg	aaacagc	cagcag	agctgtt	4860
1813	tcgatcg	aaa	ccccggc	gaaatca	ccccc	actt	4920
1814	aacattcc	tagacac	agt	atctca	ttcg	ttgaa	4980

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1817	atatctagct	aacttggct	tttcgttgt	ttttttttac	tcctttccct	cactttctcc	5160
1818	agtgcgtcaac	tgttagatat	taatcttggc	aaactgttta	atcttggta	ttttgttagat	5220
1819	ggttcaaat	gactgaactg	cattcagatt	tacagtgaa	aggaaaaatt	gcattagttg	5280
1820	gttgcgtt	cttcaagggg	cagatattac	tgcacaaact	gccatctcg	ttcatttttt	5340
1821	taactatgca	tttggataca	gactaatttt	taaaatatgc	taaactggaa	gattaaacag	5400
1822	atgtggccca	aactgttctg	gatcaggaaa	gtcatactgt	tcactttcaa	gttggctgtc	5460
1823	ccccccgcg	ccccccccc	cccccatatg	tacagatgt	aatagggtgt	ggaatgtcg	5520
1824	cagtggcaaa	cattcacag	atttttttt	tgtttctgtc	ttcaacattt	ttgacactgt	5580
1825	gctaatagtt	atattcagta	catgaaaaga	tactactgt	ttgaaagcct	tttaggaaat	5640
1826	tttgcacaga	ttttgtaca	aaacattttt	ttgaaaaat	acttgttaat	ttattctatt	5700
1827	ttaatttgc	aatgtcaata	aaaagttaag	aaataactt	ttttctagaa	gtcatttggg	5760
1828	ggtggttgtt	ccctttgtt	gttttttcc	ccccgtctt	gagttgaaca	ctattgtga	5820
1829	gagtaagcat	tccaaaggat	aaattacagg	acactaaaac	aggtcatgt	gagcttaagc	5880
1830	ggagagcagg	attnaacata	attggcataa	tgcttcattt	ttatcattgt	aacatgcctc	5940
1831	ttgggtgtct	ttaatcaaaa	gctgcaaagt	tgtcactgt	ttttttttt	tcttaattgc	6000
1832	catcatatca	agtgtactcc	agagtttagaa	aggtttgcaa	tactcaacat	tatcttttc	6060
1833	aatgggcagg	aggcaaaaaaa	aatcaagtgt	ttctgttat	acctgattca	actactaaa	6120
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E--> 1835

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1838 <211> LENGTH: 1919

1839 <212> TYPE: DNA

1840 <213> ORGANISM: Homo sapiens

1842 <220> FEATURE:

1843 <221> NAME/KEY: mRNA

1844 <222> LOCATION: (1)...(1919)

1845 <223> OTHER INFORMATION: human p35srj (MRG1)

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1851	ttcggagcag	aaatcgcaaa	aacggaaagga	ctggaaatgg	cagaccatat	gatggccatg	240
1852	aaccacgggc	gctccccga	cggcaccaat	ggctgcacc	atcacccctgc	ccaccgcatg	300
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1854	aacgcctaa	tggcgagca	catacactac	ggcgcgggca	acatgaatgc	cacgagcggc	420
1855	atcaggcatg	cgatggggcc	ggggactgt	aacggagggc	accccccgg	cgcgctggcc	480
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1857	tccctgcgg	ccagcatgca	gctgcagaag	ctcaacaacc	agtatttcaa	ccatcacccc	600
1858	taccccaaca	accactacat	gccggattt	caccctgt	caggccacca	gatgaacggg	660
1859	acaaaaccagc	actccgaga	ttgcaacccc	aagcacagcg	gccccggcag	caccccccggc	720
1860	ggctcgccgc	gcagcagcac	ccccggccgc	tctggcagca	gctcggccgg	ccggcgccggc	780
1861	agcagcaaca	gccccggcgg	cagcggcagc	ggcaacatgc	ccgcctccgt	ggcccacgtc	840
1862	cccgctgcaa	tgctgcccgc	caatgtcata	gacactgtt	tcatcgacga	ggaagtttctt	900
1863	atgtcctgg	tgtatagaaat	gggtttggac	cgcatcaagg	agctccccga	actctggctg	960
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1866	aattaaaaga	aacattccct	tagacacagt	atctcaattt	tcagatctt	aaaggtttga	1140

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1867	gaacttggaa acaaaggtaaa ctataaactt gtacaaattt gttttaaaaa aaattgctgc	1200
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1869	tgttagttgaa atatctagct aacttggtct ttttcgttgc ttgtttttac tcctttccct	1320
1870	cactttctcc agtgcctcaac tggtagatataatcttggc aaactgctta atcttggaa	1380
1871	ttttgttagat ggtttcaaat gactgaactg cattcagatt tacgagtgaa aggaaaaatt	1440
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1873	ttcattttttt taactatgca tttgagttaca gactaattttt taaaatatgc taaaactggaa	1560
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1877	acactgtgtct aatagtata ttcaagtacat gaaaagatac tactgtgttgg aaagcctttt	1800
1878	aggaaattttt gacagtattt ttgtacaaaaa catttttttgg aaaaaataact tgtaattttt	1860
1879	ttcttatttttta atttgcattt gtcataaaaaa agttaagaaa taaaaaaaaaaa aaaaaaaaaaa	1919

E--> 1884

(-31-)

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:503 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:554 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:4
L:578 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:18
L:578 M:252 E: No. of Seq. differs, <211>LENGTH:Input:5028 Found:4860 SEQ:9
L:1397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1835 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:17
L:1835 M:252 E: No. of Seq. differs, <211>LENGTH:Input:6324 Found:6180 SEQ:32
L:1884 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:1919 SEQ:33